

ANTIBODIES SPECIFIC TO HUMAN PROSTACYCLIN SYNTHASE

[01] This application is a continuation-in-part application of USSN (attorney docket Q76409) filed June 30, 2003, in turn a continuation of USSN 09/670,582 filed September 27, 2000 (now abandoned), in turn a continuation of USSN 09/037,758, filed January 10, 1998 (now abandoned), in turn a divisional of USSN 08/578,706 filed December 28, 1995 (now U.S. Patent 5,814,509)

BACKGROUND OF THE INVENTION

Technical Field

[02] The present invention relates to a polypeptide having an amino acid sequence of human-originated prostacyclin synthase (hereinafter referred to as PGIS), a DNA encoding same, a vector containing said DNA, a host cell transformed with said vector and a method for preparing human-originated PGIS comprising culturing said host cell. The present invention also relates to an antibody having a reactivity with said PGIS or its fragment. Moreover, the present invention relates to a pharmaceutical composition comprising said DNA or a vector containing said DNA, a method for promoting the production of prostaglandin I₂ and a method for the treatment of the diseases induced by a low production of prostaglandin I₂.

[03] PGIS is mainly contained in microsomal fractions of vascular endothelial cells, and is an enzyme that catalyzes synthesis of prostaglandin I₂ (hereinafter referred to as PGI₂), that is, conversion of prostaglandin H₂ (hereinafter referred to as PGH₂) to PGI₂.

[04] PGI₂ synthesized by this enzyme has potent platelet aggregation-inhibitory action and vascular smooth muscle-relaxing action. On the other hand, platelets contain thromboxane A₂ (hereinafter referred to as TXA₂) having strong platelet aggregation action and vascular

smooth muscle-contracting action, and the both substances act antagonistically in the vascular system to maintain homeostasis [British Journal of Pharmacology, vol. 76, p 3 (1982)].

[05] Cardiovascular diseases such as myocardial infarction, thrombosis and arteriosclerosis, which are among the adult diseases, have recently been considered to be caused by the imbalance in the vascular production of PGI₂ and TXA₂, particularly, insufficient vascular function due to low production of PGI₂ (ibid.).

[06] For the therapeutic treatment of the diseases presumably induced by the low production of PGI₂, PGI₂ may be supplemented as a pharmaceutical product from the outside of the body. However, PGI₂ is chemically extremely unstable to the extent that a practical use of PGI₂ itself as a pharmaceutical product may be unrealizable. In view of such situation, for example, stable PGI₂ analogs such as blood coagulation inhibitor or vasodilator are under development.

[07] The homeostasis in human and other animals which is inherently based on the balance between PGI₂ and TXA₂ may possibly destroyed by the administration of stable PGI₂ analogs. That is, administration of stable PGI₂ analogs in large amounts is associated with a risk of lowering the responsiveness of cells to PGI₂, thus impairing its capability of responding to PGI₂ when such responsiveness is in urgent need [Prostaglandins, vol. 19, p 2 (1980)].

[08] For correcting the imbalance between PGI₂ and TXA₂ and attempting the recovery of normal functions of the vascular system in an expectation of therapeutic effect over thrombosis and the like, chemically stable analogs may be used. Alongside therewith, moreover, elucidation of physicochemical property and biological property of PGIS, clarification of the relations between PGIS production and PGI₂ production while using said PGIS or DNA encoding PGIS as a research sample, and development of said PGIS or DNA encoding PGIS as pharmaceutical products to regulate the production of PGI₂ are considered

to be important and significant for the treatment of the above-said various diseases caused by the imbalance between PGI_2 and TXA_2 .

[09] Conventionally, there has been reported the tissue distribution of PGIS, namely, its presence in vascular endothelial cells, non-vascular smooth muscle cells and arterial smooth muscle of various organs [Advances in Prostaglandin, Thromboxane, and Leukotriene Research, vol. 11, pp. 87-92 (1983) and J. Biol. Chem., vol. 258, No. 9, pp. 5922-5926 (1983)]. Meanwhile, isolation and purification of PGIS from porcine and bovine have been tried [porcine: Cytochrome P450, Biochemistry, Biophysics and Environmental Implications, pp. 103-106 (1982); bovine: J. Biol. Chem., vol. 258, No. 9, pp. 3285-3293 (1983)] and N-terminal amino acid sequence and partial downstream amino acid sequence of bovine PGIS have been reported [Advances in Prostaglandin, Thromboxane, and Leukotriene Research, vol. 17, pp. 29-33 (1987) and Biochemical and Biophysical Research Communications, vol. 197, No. 3, pp. 1041-1048 (1993)].

[10] However, isolation, purification and amino acid sequence of human PGIS have not been elucidated.

DISCLOSURE OF THE INVENTION

[11] An object of the present invention is to clarify an amino acid sequence of PGIS derived from human and provide said human-originated PGIS and DNA encoding said PGIS.

[12] Said PGIS and DNA encoding said PGIS are useful as reagents for (1) the analysis of the physicochemical and biological properties of PGIS at the molecular or genetic level; (2) the analysis of the mechanism controlling the production of PGIS and the mechanism controlling the production of PGI_2 by PGIS; and (3) the investigation of the cause of various cardiovascular diseases considered to be induced by the production imbalance between PGI_2 and TXA_2 , and the molecular or genetic level analysis for the development of therapeutic

agents for said diseases. In addition, PGIS and its mRNA are useful as diagnostics for the determination of expression level and distribution in the body tissues. Still further, they are expected to provide therapeutic agents for, for example, various cardiovascular disorders such as thrombosis, myocardial infarction, arteriosclerosis and angina pectoris, which enhance the production level of PGI₂ upon introduction of themselves, fragment thereof or modified compound thereof into the body in a lesion-specific manner.

[13] Another object of the present invention is to provide a recombinant vector containing a DNA encoding human-originated PGIS, the expression system of PGIS which comprises a host cell transformed with said vector, and a method for preparing PGIS by genetic engineering using said expression system.

[14] According to such method, human-originated PGIS can be prepared in great amounts with ease and with high efficiency.

[15] The present invention also aims at providing a human-originated PGIS antibody useful for the purification of human-originated PGIS and immunohistochemical analysis of the cause of a disease.

[16] The present inventor has conducted intensive studies with the aim of accomplishing the above-mentioned objects, and succeeded in cloning cDNA encoding PGIS from human aorta endothelial cells and identifying the primary structure of human-originated PGIS from the nucleotide sequence of said cDNA, which resulted in the completion of the present invention.

[17] Accordingly, the present invention relates to a DNA comprising a DNA having a nucleotide sequence encoding an amino acid sequence of human-originated PGIS substantially depicted in SEQ ID NO: 15, preferably a DNA comprising a DNA having a 28th-1527th nucleotide sequence substantially shown in SEQ ID NO: 14, and more preferably a DNA having a 28th-1527th nucleotide sequence shown in SEQ ID NO: 14.

[18] The present invention also relates to a recombinant vector comprising the above-mentioned DNA, a host cell transformed with said vector and a method for preparing human-originated PGIS comprising culturing said host cell in a medium and recovering human-originated PGIS from the obtained culture.

[19] The present invention also relates to a polypeptide having an amino acid sequence of human-originated PGIS which is substantially shown in SEQ ID NO: 15, and antibodies having reactivities with said human-originated PGIS.

[20] The present invention further relates to a pharmaceutical composition comprising said DNA or a recombinant vector comprising said DNA. Said pharmaceutical composition can be used as a medicament for promoting PGI₂ production or for treating the diseases induced by a low production of PGI₂.

[21] The present invention moreover relates to a method for promoting the production of PGI₂, comprising introducing the above-mentioned DNA or a recombinant vector comprising said DNA into human or other animals. The present invention also relates to a method for treating the diseases induced by a low production of PGI₂, comprising introducing the above-mentioned DNA or a recombinant vector comprising said DNA into human or other animals.

BRIEF DESCRIPTION OF THE DRAWINGS

[22] FIG. 1 shows a restriction enzyme map of human PGIS cDNA, and the PGIS DNA region comprised in λ hPGIS141, pHPGIS135 and pHPGIS36.

[23] FIG. 2 shows a restriction enzyme map of plasmid pHPGIS36.

[24] FIG. 3 shows a restriction enzyme map of plasmid pHPGIS135.

[25] FIG. 4 shows a restriction enzyme map of plasmid pHPGIS1.

[26] FIG. 5 shows a restriction enzyme map of human PGIS expression vector pCMV-HPGIS1.

- [27] FIG. 6 shows an expression vector pUC-CAGGS.
- [28] FIG. 7 is a photograph showing the results of the analysis, by thin layer chromatography, of the PGIS activity in the cells into which pCMV-HPGIS1 has been introduced.
- [29] FIG. 8 is a photograph showing the results of the analysis, by thin layer chromatography, of the PGIS activity in positive control (bovine platelet microsomes).
- [30] FIG. 9 is a photograph showing the results of the analysis, by thin layer chromatography, of negative control wherein pCMV alone was introduced.
- [31] FIG. 10 is a graph showing the effects of the introduction of human PGIS expression vector on the blood vessel smooth muscle cell proliferation.
- [32] FIG. 11 is a photograph showing the results of RNA blot (electrophoresis) analysis of human PGIS mRNA treated with cytokines.
- [33] FIG. 12 is a photograph showing the distribution of PGIS mRNA expression in human body (pancreas, kidney, skeletal muscle, liver, lung, placenta, brain and heart) by electrophoresis.
- [34] FIG. 13 is a photograph showing the distribution of PGIS mRNA expression in human body (peripheral leukocyte, large intestine, small intestine, ovary, testicle, prostate, thymus and spleen) by electrophoresis.
- [35] FIG. 14 depicts a Western blot analysis of Example 7.

DETAILED DESCRIPTION OF THE INVENTION

- [36] The present invention is explained in detail in the following.
- [37] The polypeptide of the present invention has a catalytic activity to convert PGH_2 to PGI_2 and has an amino acid sequence of human-originated PGIS substantially shown in Sequence Listing, SEQ ID NO: 15 to be mentioned later.

[38] By "substantially" is meant that the polypeptide of the present invention is not limited to the polypeptide having the amino acid sequence shown in SEQ ID NO: 15, but may include deletion, substitution and addition with respect to some of the amino acids in the amino acid sequence shown in SEQ ID NO: 15, as long as the polypeptide has immunological and biological activity (human PGIS activity) similar to that of human-originated PGIS having said amino acid sequence.

[39] While the site of deletion, substitution and addition of the amino acids is not particularly limited, at least 441st Cys residue and thereabout region in the amino acid sequence shown in Sequence No. 15 need to be reserved. This is because human-originated PGIS of the present invention is homologous to known cytochrome P450 in the amino acid sequence, since it has Cys residue in the C-terminal side of the amino acid sequence constituting the heme-binding site (fifth ligand) which is important for the expression of biological activity of cytochrome P450, and speculated to be a new protein belonging to the cytochrome P450 family [see Seibutsu Butsuri, vol. 32, No. 1, pp. 10-15 (1992)].

[40] The polypeptide of the present invention preferably has an amino acid sequence of human-originated PGIS shown in SEQ ID NO: 15.

[41] The PGIS activity possessed by the polypeptide of the present invention is a catalytic activity to convert PGH_2 to PGI_2 . Said PGIS activity can be determined according to the method of Salmon, J. A. and Flower, R. J. et al [Methods Enzymol., 86, pp. 91-99 (1982)] wherein the conversion of ^{14}C -labeled PGH_2 to PGI_2 is assayed by separating the metabolite of 6-keto- $\text{PGF}_{1\alpha}$ by thin layer chromatography and detecting the radioactivity of said 6-keto- $\text{PGF}_{1\alpha}$.

[42] The present invention also relates to a DNA comprising a DNA having a nucleotide sequence encoding the amino acid sequence of human-originated PGIS substantially shown in SEQ ID NO: 15.

[43] Said DNA may be any as long as it comprises a DNA having a nucleotide sequence encoding the aforementioned amino acid sequence of human-originated PGIS, and is exemplified by a DNA encoding the polypeptide having the amino acid sequence shown in SEQ ID NO: 15 or a polypeptide having the equivalent immunological and biological activity. More specifically, it is a DNA comprising the 28th-1527th nucleotide sequence in the nucleotide sequence shown in SEQ ID NO: 14.

[44] In general terms, the genetic recombinant technique enables conversion of at least one nucleotide of a DNA sequence of a gene to a different nucleotide according to the degeneracy of the genetic code, without changing the amino acid sequence of a protein produced by the gene. Accordingly, the DNA of the present invention encompasses a DNA comprising a nucleotide sequence obtained by modification for substitution, based on the genetic code, of the 28th-1527th nucleotide sequence of Sequence Listing SEQ ID NO: 14.

[45] The DNA of the present invention can be obtained by any method. For example, the present invention encompasses complementary DNA (cDNA) prepared from mRNA, DNA prepared from genomic DNA, DNA obtained by chemical synthesis, DNA obtained by amplification by PCR using RNA or DNA as a template, and DNA constructed by suitably combining these methods.

[46] The DNA of the present invention can be obtained by a method comprising cloning cDNA from mRNA of human-originated PGIS by a conventional method, a method comprising splicing an isolated genomic DNA for PGIS, a method comprising chemical synthesis or other method.

[47] (1) For example, a method for cloning cDNA from mRNA encoding human-originated PGIS comprises the following steps.

[48] Cells producing human-originated PGIS, such as human aorta endothelial cells are cultured and mRNA encoding said PGIS is prepared from the culture thereof. mRNA is

prepared by, for example, applying entire RNA prepared by a known method such as guanidine thiocyanate method [Chirgwin, J. M. et al., *Biochem.*, 18, 5294 (1979)], heat phenol method and AGPC to affinity chromatography using oligo(dT)-cellulose or poly U-sepharose.

[49] Using the obtained mRNA as a template, cDNA chain is synthesized by a known method using a reverse transcriptase [e.g., the method of Okayama, H. et al: Okayama, H. et al., *Mol. Cell. Biol.*, 2, 161 (1982) and *ibid.* 3, 280 (1983), and the method of Gubler, U. and Hoffman, B. J.: Gubler, H. and Hoffman, B. J., *Gene*, 25, 263 (1983)], thereby converting the same to a double stranded cDNA. This cDNA is inserted into a plasmid vector or a phage vector, with which *Escherichia coli* is transformed, or transfected after in vitro packaging, to prepare cDNA library.

[50] The plasmid vector used here is not subject to any particular limitation as long as it can be retained by replication in the host, and the phage vector is not limited either as long as it can proliferate in the host. Examples of the conventionally-used cloning vector include pUC119, λ gt10 and λ gt11. When immunological screening to be mentioned later is to be employed, the vector preferably contains a promoter capable of expressing the PGIS gene in the host.

[51] The method for insertion of a cDNA into plasmid is exemplified by a method described in Maniatis, T. et al [*Molecular Cloning, A Laboratory Manual*, Cold Spring Harbor Laboratory, p. 239 (1982)]. The method for insertion of a cDNA into phage vector includes the method of Hyunh, T. V. et al [*DNA Cloning, a practical approach*, 1, 49 (1985)]. For simplification, a commercially available ligation kit (e.g., those manufactured by Takara Shuzo) can be used. The recombinant plasmid and phage vector thus obtained are introduced into a suitable host such as prokaryotic cells (e.g., *E. coli* HB101, DH5 and MC1061/P3).

[52] The method for introducing a plasmid into a host includes calcium chloride method and calcium chloride/rubidium chloride method described in Maniatis, T. et al [Molecular Cloning, A Laboratory Manual, Cold Spring Harbor Laboratory, p. 239 (1982)] and electroporation method. The method for introducing a phage vector into a host is exemplified by a method comprising in vitro packaging of phage DNA and introducing same into proliferated host cells. In vitro packaging can be carried out easily by using a commercially available in vitro packaging kit (e.g., product of Stratagene and product of Amersham).

[53] The cDNA encoding the PGIS of the present invention can be isolated from the cDNA library prepared by the above method, by a combination of general cDNA screening methods.

[54] Such methods include, for example, a method wherein an oligonucleotide considered to be corresponding to the amino acid sequence of human PGIS is chemically synthesized separately and labeled with ^{32}P to give a probe, and a clone having the desired cDNA is screened by a known colony hybridization [Crunstein, M. and Hogness, D. S., Proc. Natl. Acad. Sci. USA, 72, 3961 (1975)] or plaque hybridization [Molecular Cloning, A Laboratory Manual, Cold Spring Harbor Laboratory, p. 239 (1982)]; and a method wherein PCR primer is prepared and a specific region of PGIS is amplified by PCR method, which is followed by selecting a clone having a DNA fragment encoding said region. When a cDNA library prepared using a vector (e.g. λ gt11 phage vector) capable of expressing cDNA is used, the objective clone can be selected based on an antigen-antibody reaction using the PGIS antibody of the present invention to be mentioned later. When large amounts of clone are treated, screening based on PCR is preferable.

[55] The nucleotide sequence of DNA thus obtained can be determined by Maxam-Gilbert method [Maxam, A. M. and Gilbert, W., Proc. Natl. Acad. Sci. USA., 74, 560 (1977)] or synthetic dideoxynucleotide chain termination method using phage M13 [Sanger, F. et al,

Proc. Natl. Acad. Sci. USA., 74, 5463-5467 (1977)]. The PGIS gene can be obtained by cleaving all or part thereof from the clone obtained above by using a restriction enzyme and the like.

[56] (2) A preparation method comprising isolating DNA encoding PGIS from genomic DNA of human aorta vascular cells includes, for example, the following method.

[57] Human aorta vascular cells are lysed preferably using SDS or proteinase K, and DNA is deproteinized by repetitive extraction with phenol. RNA is preferably digested with ribonuclease. The obtained DNA is partially digested with a suitable restriction enzyme and the obtained DNA fragment is amplified by a suitable phage or cosmid to form a library. Then, the clone having the desired sequence is detected by, for example, a method using a DNA probe with a radioactive label, and a whole or partial PGIS gene is cleaved from said clone by using a restriction enzyme and the like.

[58] (3) The DNA of the present invention can be prepared by chemical synthesis by a conventional method based on the nucleotide sequence depicted in Sequence Listing SEQ ID NO: 14.

[59] The present invention further relates to a recombinant vector comprising DNA encoding the above-mentioned PGIS. The recombinant vector of the present invention is not particularly limited as long as it can be retained by replication or self-proliferation in various prokaryotic and/or eukaryotic host cells, and includes plasmid vector and phage vector.

[60] The recombinant vector can be easily prepared by ligating the DNA encoding human-originated PGIS of the present invention with a commercially available recombinant vector (plasmid DNA and bacteriophage DNA) by a conventional method. Usable recombinant vector includes, for example, Escherichia coli-originated plasmids pBR322, pBR325, pUC12 and pUC13; yeast-originated plasmids pSH19 and pSH15; and Bacillus subtilis-originated plasmids pUB110, pTP5 and pC194. Examples of phage include bacteriophage such as λ

phage, and animal or insect viruses such as retrovirus, vaccinia virus, nuclear polyhedrosis virus and adenovirus [e.g. pVL1392, pBK283, Autographa californica nuclear polyhedrosis virus (AcNPV) and Bombyx mori nuclear polyhedrosis virus (BmNPV)].

[61] When production of PGIS by the expression of the PGIS gene is aimed, an expression vector is useful. The expression vector is not particularly limited as long as it expresses the PGIS gene in various prokaryotic and/or eukaryotic host cells and is capable of producing proteins. Preferred are that derived from insect virus which infects insect cells and produces PGIS in said cells, and that derived from animal virus which infects animal cells and produces PGIS in said cells.

[62] When bacteria, particularly Escherichia coli, is used as the host cell, the expression vector generally consists of at least promoter-operator region, initiation codon, DNA encoding the PGIS of the present invention, termination codon, terminator region and replicon.

[63] When yeast, animal cell or insect cell is used as the host cell, the expression vector preferably consists of at least promoter, initiation codon, DNA encoding the polypeptide of the present invention and termination codon. It may contain DNA encoding signal peptide, enhancer sequence, non-translation region on the 5' or 3' side of the polypeptide of the present invention, splicing junction, polyadenylation site, selection marker region, replicon and the like.

[64] The promoter-operator region for expressing the polypeptide of the present invention in bacteria contains promoter, operator and Shine-Dalgarno (SD) sequence such as AAGG. When the host is Escherichia coli, the region preferably contains, for example, Trp promoter, lac promoter, recA promoter, λ PL promoter and lpp promoter. The promoter for expressing PGIS in yeast includes, for example, PH05 promoter, PGK promoter, GAP promoter and ADH promoter, and when the host is bacteria belonging to the genus Bacillus, SL01

promoter, SP02 promoter and penP promoter can be used. When the host is eukaryotic cells such as animal cells, examples of the promoter include, but not limited to, SV40-derived promoter, retrovirus promoter, heat shock promoter, polyhedron promoter that a nuclear polyhedrosis virus has, cytomegalovirus promoter, adenovirus promoter and β -actin promoter. The use of an enhancer is also effective for the expression.

[65] Preferable initiation codon includes, for example, methionine codon (ATG).

[66] The termination codon is exemplified by conventional termination codons such as TAG and TGA.

[67] As the terminator region, conventional intact or synthetic terminator can be used.

[68] By replicon is meant a DNA capable of reproducing the entire DNA sequence in the host cell, and exemplified by naturally occurring plasmid, artificially modified plasmid (DNA fragment prepared from naturally occurring plasmid) and synthetic plasmid. Examples of preferable plasmid include plasmid pBR322 and artificial modification thereof (DNA fragment obtained by treating pBR322 with a suitable restriction enzyme) in the case of *E. coli*; yeast 2 μ plasmid and yeast chromosomal DNA in the case of yeast; and plasmid pRSVneo ATCC 37198, plasmid pSV2dhfr ATCC 37145, plasmid pdBPV-MMTneo ATCC 37224 and plasmid pSV2neo ATCC 37149 in the case of mammalian cell.

[69] Enhancer sequence, polyadenylation site and splicing junction site can be those conventionally used by artisan, such as respective ones derived from SV40.

[70] As the selection marker, conventional ones can be used according to a conventional method. Examples thereof include a gene resistant to antibiotic such as tetracycline, ampicillin and kanamycin.

[71] The expression vector of the present invention can be prepared by ligating at least the above-mentioned promoter, initiation codon, DNA encoding PGIS of the present invention, termination codon and terminator region sequentially and cyclically into a suitable

replicable unit. For this end, suitable DNA fragments such as linker and other restriction sites can be used by a conventional method such as digestion with restriction enzyme and ligation using T4DNA ligase on demand.

[72] The transformant of the present invention can be prepared by introducing the above-mentioned expression vector into a host cell.

[73] Examples of the host cell include microorganisms such as bacteria (e.g. bacteria belonging to the genera *Escherichia* and *Bacillus*), yeast such as those belonging to the genus *Saccharomyces*, animal cells and insect cells. Specifically exemplified are *Escherichia coli* K12DH1, M103, JA221, HB101, C600, XL-1 Blue and JM109 as the bacteria belonging to the genus *Escherichia*; and *Bacillus subtilis* 207-21 as the bacteria belonging to the genus *Bacillus*. Examples of the yeast include *Saccharomyces cerevisiae* AH22, AH22R-, NA87-11A and DKD-5D. Examples of animal cell include simian cell COS-7, Vero, Chinese hamster cell CHO, mouse L cell, human FL cell and human 293 cell. Examples of insect cell include BmN4 and Sf9. Preferred are insect cells and animal cells.

[74] The preferred host cell for cloning the DNA sequence and constructing the vector is generally a prokaryotic cell. The expression vector constructed is used to transform a suitable host cell. The host cell may be a prokaryotic cell or an eukaryotic cell as well. Preferred are insect cells (e.g., BmN4 and Sf) and animal cells.

[75] The expression vector is introduced (i.e., transformation which is used in a concept inclusive of transfection in the present invention) into host cells by a conventionally known method.

[76] For example, in the case of bacteria (e.g. *Escherichia coli* and *Bacillus subtilis*), the method of Cohen et al [Proc. Natl. Acad. Sci. USA., 69, 2110 (1972)] protoplast method [Mol. Gen. Genet., 168, 111 (1979)] or competent method [J. Mol. Biol., 56, 209 (1971)] may be used; in the case of *Saccharomyces cerevisiae*, the method of Hinnen et al [Proc.

Natl. Acad. Sci. USA., 75, 1927 (1978)] or lithium method [J. Bacteriol., 153, 163 (1983)] may be used; in the case of animal cells, the method of Graham [Virology., 52, 456 (1973)], lipofectin method or HVJ-liposome method [Hypertension, 21, 894-899 (1993)] may be used; and in the case of insect cells, the method of Summers et al [Mol. Cell. Biol., 3, 2156-2165 (1983)] may be used for transformation.

[77] The human-originated PGIS of the present invention can be prepared by culturing, in a nutrient medium, a transformant (which term is used in a concept inclusive of transfectant in the present invention) comprising the expression vector prepared as in the above.

[78] The nutrient medium preferably contains carbon source, inorganic nitrogen source or organic nitrogen source necessary for the growth of host cell (transformant). Examples of carbon source include glucose, dextran, soluble starch and sucrose; examples of inorganic nitrogen source or organic nitrogen source include ammonium salts, nitric acid salts, amino acid, corn steep liquor, peptone, casein, meat extract, soybean meal and potato liquid extract. When desired, other nutrients such as inorganic salt (e.g. calcium chloride, sodium dihydrogenphosphate and magnesium chloride), vitamins, and antibiotics such as ampicillin and kanamycin may be added to the medium.

[79] Culture is carried out according to the method known in the pertinent field. Culture conditions such as temperature, pH of the medium and culture time are appropriately determined so that the maximum potency of PGIS can be obtained.

[80] Specific media and culture conditions to be employed according to the host cell are exemplified in the following, which are not limitative.

[81] When the host is bacteria, Actinomyces, yeast or filamentous fungus, for example, liquid media containing the above-mentioned nutrient sources are appropriate. Preferred is a medium having a pH of 5-8.

- [82] When the host is *Escherichia coli*, preferable medium is M9 medium [Miller, J., Exp. Mol. Genet., p. 431, Cold Spring Harbor Laboratory, New York (1972)]. In this case, culture is performed with aeration and agitation as necessary, at 14°C-43°C for about 3 to 24 hours.
- [83] When the host is bacteria belonging to the genus *Bacillus*, culture is performed with aeration and agitation as necessary, at 30°-40°C for about 16 to 96 hours.
- [84] When the host is yeast, the medium is exemplified by Burkholder minimum medium [Bostian, K. L. et al, Proc. Natl. Acad. Sci. USA, 77, 4505 (1980)] which preferably has a pH of 5-8. Culture is generally performed at about 20-35°C for about 14 to 144 hours with aeration and agitation where necessary.
- [85] When the host is animal cell, the medium is exemplified by MEM medium containing fetal calf serum at about 5-20% [Science, 122, 501 (1952)], DMEM medium [Virology, 8, 396 (1959)], RPMI1640 medium [J. Am. Med. Assoc., 199, 519 (1967)] and 199 medium [Proc. Soc. Exp. Biol. Med., 73, 1 (1950)]. The pH of the medium is preferably about 6-8, and culture is generally performed at about 30°C-40°C for about 15-60 hours with aeration and agitation where necessary.
- [86] When the host is insect cell, the medium is exemplified by Grace's medium containing fetal calf serum [Proc. Natl. Acad. Sci. USA, 82, 8404 (1985)] which preferably has a pH of about 5-8. Culture is generally performed at about 20°C-40°C. for about 15 to 100 hours with aeration and agitation where necessary.
- [87] The human-originated PGIS of the present invention can be recovered as in the following from the culture obtained above.
- [88] That is, when the human-originated PGIS is present in the liquid portion of the culture, the culture thus obtained is subjected to filtration or centrifugation to separate culture

filtrate (supernatant), and PGIS is purified and separated from said culture filtrate by a conventional method employed for purifying and isolating natural or synthetic proteins.

[89] The method for purification and isolation includes, for example, a method utilizing the solubility, such as salting out and solvent precipitation, a method utilizing the difference in molecular weights such as dialysis, ultrafiltration, gel filtration and sodium dodecyl sulfate-polyacrylamide gel electrophoresis, a method utilizing charge such as ion exchange chromatography and hydroxyapatite chromatography, a method utilizing specific affinity such as affinity chromatography, a method utilizing the difference in hydrophobicity such as reversed phase high performance liquid chromatography and a method utilizing difference in isoelectric point such as isoelectric focusing.

[90] When the human-originated PGIS is present in the periplasm or cytoplasm of the cultured transformant, the culture is subjected to a conventional method such as filtration and centrifugation to collect the cells; the cells are suspended in a suitable buffer and subjected to lysis of cell wall and/or cell membrane by ultrasonication, using lysozyme or by freeze-thawing; and the membrane fraction containing PGIS is obtained by centrifugation or filtration. Said membrane fraction is solubilized with surfactant such as Triton to give a crude solution. The crude solution is treated by a conventional method as exemplified supra to isolate and purify PGIS of the present invention.

[91] The present invention also relates to an antibody having a reactivity with the above-said human-originated PGIS. The antibody of the present invention encompasses both the polyclonal antibody and monoclonal antibody having the above-mentioned properties. The antibody of the present invention can be obtained by a conventional method.

[92] For example, the monoclonal antibody of the present invention can be prepared from hybridoma produced by so-called cell fusion. That is, fused hybridoma is formed from the antibody-producing cell and bone marrow cell; said hybridoma is cloned; and a clone is

selected which produces an antibody having a specific affinity for an antigen, i.e. a polypeptide having part or whole of the human-originated PGIS amino acid sequence. The procedure therefor may be known methods except the use of the human-originated PGIS of the present invention as an immunizing antigen.

[93] The immunogen can be used for immunizing animals after admixing with, for example, complete Freund adjuvant. Examples of the animal include mouse, rat and rabbit. The animals are immunized by subcutaneous, intramuscular or intraperitoneal injection of about 5-200 μg /injection. The immunization includes 1-4 times of immunization at about every 1-2 weeks from the initial immunization and final immunization at about 1-4 weeks thereafter. When about 3-5 days have passed since final immunization, antibody-producing cells are separated from the immunized animal. The antibody-producing cells are exemplified by spleen cells and lymph node cells.

[94] The bone marrow cells are, for example, those derived from mouse, rat and human. Examples thereof include mouse myeloma P3.X63.Ag8, P3.X63.Ag8-U1, P3.NS1-Ag4, SP2/0-Ag14 and X63-Ag8 . 653. It is preferable that the antibody-producing cells and bone marrow cells be derived from the same species of animals.

[95] Cell fusion is performed by the method described in, for example, Nature, vol. 266, p. 550 (1977) or an analogous method. Specifically, it is performed using 30-50% polyethylene glycol having an average molecular weight of 1,000-4,000 at 30°C-40°C for about 1-3 minutes.

[96] The cells obtained by cell fusion are subjected to screening for a clone which produces the desired monoclonal antibody. That is, the cells are cultured in, for example, a microplate and the antibody titer of the culture supernatant in the well in which cell growth was acknowledged is determined by, for example, enzyme antibody method to obtain the well in which suitable antibody has been produced. Cloning by, for example, limiting dilution

from such well gives clones. The monoclonal antibody of the present invention can be obtained by culturing said hybridoma cell clone by conventional culture method, high density culture method or spinner-flask culture method and purification thereof by affinity chromatography using protein A-bound carrier or anti-mouse immunoglobulin-bound carrier.

[97] Alternatively, the cultured hybridoma cells are intraperitoneally injected to the mouse of the same species which has been previously treated with pristane, and ascites obtained is subjected to salting out with ammonium sulfate and DEAE ion exchange chromatography to give purified IgG fraction containing the same.

[98] The DNA encoding the human-originated PGIS of the present invention can be used for gene therapy.

[99] The DNA encoding the human-originated PGIS of the present invention or a recombinant vector comprising said DNA is introduced into human or other animals, whereby PGIS is produced in the human or other animals to promote production of PGI₂. The promoted PGI₂ production in turn enables treatment (therapeutic treatment or improvement of symptoms) of the diseases induced by a low production of PGI₂. Examples of the diseases induced by the low production of PGI₂ include cardiovascular diseases such as thrombosis, myocardial infarction, arteriosclerosis and angina pectoris. The recombinant vector may be introduced into human or other animals in the form of cells transformed with said recombinant vector.

[100] The gene therapy utilizing the gene (inclusive of DNA and recombinant vector) of the present invention permits setting an appropriate environment in which the gene of the present invention introduced into a human or other animal can fully show its function. The treatment can be given by a conventional method as long as it intends expression of desired effects of the gene of the present invention. Such method is exemplified by virological means utilizing retrovirus vector or adenovirus vector, physical means for introducing gene by particle gun

method or by using naked DNA, and chemical means such as lipid method [Molecular Medicine, vol. 30, No. 12, p. 1526 (1993); Jikken Igaku, vol. 12, No. 3, p. 15, 28 and 40 (1994); Proc. Natl. Acad. Sci. USA, 92, 1137 (1995)]. A method using an adenovirus vector which can be used for the gene therapy of cystic fibrosis and which is known to permit efficient introduction of gene into differentiated cells and tissues and expression therein, and a method using a fusogenic liposome which allows introduction of optional gene into tissue cells in vivo are preferable for the gene treatment of the present invention.

[101] The dose of the DNA or recombinant vector of the present invention is subject to appropriate change according to sex, age and body weight of patients, the kind of disease and symptoms thereof, and administration route. For example, 100 µg-10 mg of DNA is generally administered.

[102] The DNA and recombinant vector of the present invention are administered by intravenous injection, transmucosal administration, oral administration using enteric-coated agents, or topical administration, with preference given to topical administration using catheter and the like.

[103] The DNA encoding human-originated PGIS and recombinant vector comprising said DNA of the present invention are admixed with conventional, pharmaceutically acceptable carrier, excipient, diluent, extender, disintegrator, stabilizer, preservative, buffer, emulsifier, flavor, coloring, sweetener, thickener, elixir, solubilizer and other additives such as water, salt solution, phosphate buffer, vegetable oil, ethanol, polyethylene glycol, glycerol, gelatin, lactose, glucose, mannitol, starch, sucrose, magnesium stearate, hydroxypropylcellulose, talc, lanolin and petrolatum, and can be used in the form of injection, tablet, powder, capsule, enteric-coated agent, ointment, suspension, emulsion, spray, inhalant, collunarium and the like.

[104] A pharmaceutical composition comprising the DNA or recombinant vector comprising said DNA of the present invention can be administered to mammals such as human, mouse, rat, rabbit, pig, cow, sheep, dog and cat.

EFFECTS OF THE INVENTION

[105] The present invention gives the first clarification of the amino acid sequence of human-originated PGIS and nucleotide sequence of DNA encoding the enzyme having said sequence. Based on the elucidation of such amino acid sequence and nucleotide sequence, the present invention provides a method for preparing PGIS by genetic engineering and an expression system related thereto.

[106] The PGIS and DNA encoding same of the present invention are useful as reagents for

- (1) the analysis of physicochemical property and biological property of PGIS at the molecular or genetic level,
- (2) the analysis of the mechanism of regulating PGIS production and the mechanism of regulating PGI₂ production by PGIS, and
- (3) the investigation of the cause of various cardiovascular diseases considered to be induced by the production imbalance between PGI₂ and TXA₂, and analysis at the molecular or genetic level for the development of therapeutic agent for said diseases.

[107] In addition, they are useful as diagnostics for determining the in vivo tissue expression level and distribution of PGIS or mRNA thereof.

[108] Moreover, they can be used as therapeutic agents for various cardiovascular diseases such as thrombosis, myocardial infarction, arteriosclerosis and angina pectoris, which

increase the production level of PGI₂ based on lesion-specific introduction, into human and other animals, of PGIS, DNA encoding PGIS, fragment thereof or modified product thereof.

[109] The expression system of PGIS comprising a recombinant vector containing DNA encoding the human-originated PGIS of the present invention, and a host cell transformed with said vector is useful for the production by genetic engineering, which enables easy and efficient mass production of human-originated PGIS.

[110] In addition, the human-originated PGIS antibody of the present invention serves well for the purification of human-originated PGIS and the immunohistochemical analysis of the cause of a disease (specific staining of various tissues such as uterus, heart, skeletal muscle, lung and prostate).

[111] The plasmid, enzyme such as restriction enzyme, T4DNA ligase and other substances to be used in Examples of the present invention are commercially available and can be used according to a conventional method. The procedures for cloning of cDNA, determination of nucleotide sequence, transfection of host cell, culture of transfectant, harvesting and purification of PGIS from obtained culture, and obtainment of antibody are well known to those skilled in the art, or can be known from literatures.

[112] The pHPGIS36 (PBJT-BA-4, deposit number FERM BP-4653) and pHPGIS135 (PBJT-BA 5, deposit number FERM BP-4654) used in the present invention are at international deposit at National Institute of Bioscience and Human-Technology Agency of Industrial Science and Technology.

EXAMPLES AND REFERENCE EXAMPLES

The present invention is described detailedly in the following by way of Examples and Reference Examples, to which the present invention is not limited.

Example 1

Determination of cDNA nucleotide sequence

(1) Preparation of λ hPGIS141

[113] Human genomic library (Genomic lung fibroblast cell line, WI 38, manufactured by Clone-Tech) was seeded at about 2×10^5 PFU, and screened by plaque hybridization using, as a probe, a bovine cDNA prepared in advance by the inventor (see Tanabe, T., Hara, S., Miyata, A., Brugger, R., and Ullrich, V. (1993) in Abstract book of 3rd international conference on eicosanoid and other bioactive lipids in cancer, inflammation and radiation Injury, p. 137).

[114] As a result, four positive signals were obtained, one of which was isolated to a single plaque. Liquid culture thereof resulted in mass preparation of phage DNA. After purification, it was digested with various restriction enzymes, followed by mapping. A fragment comprising exon was identified by Southern hybridization, which was followed by structural analysis by DNA sequencing to confirm that the finally-isolated clone (λ hPGIS141) coded for human PGIS.

[115] The λ hPGIS141 thus obtained was structurally analyzed by restriction enzyme site mapping and nucleotide sequence determination, and it was found that λ hPGIS141 contained the region corresponding to a 673rd-855th nucleotide sequence of bovine PGIS cDNA (SEQ ID NO: 8) corresponding to amino acid SEQ ID NO: 9.

[116] Based on the nucleotide sequence of λ hPGIS141 cDNA fragment thus obtained, primers [SEQ ID NO: 1: P1 primer (674-689), SEQ ID NO: 2: P2 primer (699-718), SEQ ID

NO: 3: P3 primer (696-713), SEQ ID NO: 4: P4 primer (805-822)] having the sequences depicted in Sequence List SEQ ID NOS: 1-4 were synthesized.

(2) Amplification of cDNA by PCR method

[117] The 3'-downstream region and 5'-upstream region of cDNA were amplified by PCR method [Biochem. Biophys. Res. Commun. 178, 1479-1484 (1991)] using said primers and poly(A)⁺ RNA (mRNA) from 1 µg of human aorta vascular endothelial cells (hereinafter referred to as HAEC, manufactured by Kurabo) as a template.

[118] For amplification of cDNA corresponding to the 3'-downstream region, cDNA was primed with a dT₁₇ adapter (5'-GACTCGAGTCGACATCGA-(T)₁₇-3', SEQ ID NO: 5), and elongated to give a first cDNA chain which was amplified with P1 primer (674-689) and the adapter primer (SEQ ID NO: 6), and then with P2 primer (699-718) and the adapter primer (SEQ ID NO: 6). The 5'-upstream region of the cDNA was amplified using a 5' RACE system (GIBCO BRL). According to the protocol, homomeric dC tail was added to the first cDNA chain and a second cDNA chain was formed using an adapter primer (5'-(CUA)₄ GGCCACGCGTCGACTAGTACGGGIIIGGGIIIGGGIIG-3') (SEQ ID NO: 7). The first step amplification was performed using P4 primer and the adapter primer (SEQ ID NO: 7). The second step amplification was performed using P3 primer and the adapter primer (SEQ ID NO: 7). The PCR method was repeated 35 cycles according to the following cycloprofile.

Denaturation	94°C, 1 minute
Annealing	54°C, 1 minute
Elongation	72°C, 3 minutes

[119] The respective PCR products (3'-downstream region amplification product and 5'-downstream region amplification product) were partially taken out and purified by

electrophoresis using 1% agarose gel. Southern hybridization was applied using bovine cDNA (pBPGISI) as a probe, and DNA was extracted from the band which cross-hybridized to said probe. The obtained DNA was cloned into pBluescriptII SK(-).

[120] That is, cloning and screening were performed by the following steps:

- (1) cleaving out the band which showed a signal from a gel, after electrophoresis
- (2) agarase digestion at 40°C for one hour (agarase 1 unit/100 µl gel)
- (3) extraction of DNA with phenol and subsequent ethanol precipitation
- (4) dissolving said DNA ethanol precipitate in sterile water and treating with polynucleotide kinase at 37°C for one hour
- (5) end repairing with Klenow fragment (16°C, 1 hr)
- (6) ligation using Takara ligation kit
- (7) transformation by a conventional method
- (8) sewing in a plate
- (9) forming a replica by a conventional method and
- (10) colony hybridization of nitrocellulose filter of the replica by a conventional method, using bovine PGIS cDNA as a probe

[121] The hybridization was performed at 60°C in 6 X SSC [1 X SSC containing 0.15M NaCl, 15 mM sodium citrate (pH 7.0)], 5 X Denhardt's solution, 250 µg/ml salmon sperm DNA, 0.1% SDS and cDNA fragment (10⁶ cpm/ml) labeled by random priming method. The filter obtained was washed twice with 3 X SSC and 0.1% SDS at room temperature for 5 minutes and twice with 0.1 X SSC and 0.1% SDS at 50°C for 15 minutes. The filter was air-dried, and exposed to Fuji X ray film using a intensifying screen at -80°C for 12-16 hr.

[122] The obtained DNA insert was subcloned into pBluescriptII SK(-). By these steps, a clone (pHPGIS135) containing 3'-downstream region DNA of human-originated PGIS and a

clone (pHPGIS36) containing 5'-upstream region DNA of human-originated PGIS were obtained. Then, the nucleotide sequence of the DNA insert of respective clones was determined by the Sanger method [Sanger, F., Nickle, S., and Coulson, A. R. (1977) Proc. Natl. Acad. Sci. USA 74, 5463-5467] using Taq dye primer cycle sequence kit (manufactured by Applied Biosystems) and Model 373A DNA sequencer (manufactured by Applied Biosystems). As a result, it was found that pHPGIS36 clone had, as a DNA insert sequence, a 740 bp nucleotide sequence (SEQ ID NO: 10) of cDNA of human PGIS, having an adapter sequence on the 5' side, based on which partial amino acid sequence of PGIS comprising 237 amino acid residues wherein ATG is the translation initiation sequence (Met) was identified. [123] It was also found that pHPGIS135 clone comprised, as a DNA insert sequence, a 1277 bp nucleotide sequence (SEQ ID NO: 12) of cDNA of human PGIS, having an adapter sequence on the 3' side, based on which partial amino acid sequence of PGIS on the carboxyl side region starting from 226th aspartic acid was identified. The nucleotide sequence of human PGIS cDNA contained in pHPGIS36 clone and the amino acid sequence deduced therefrom are depicted in SEQ ID NO: 11 in the Sequence Listing to be mentioned later, and the nucleotide sequence of human PGIS cDNA contained in pHPGIS135 clone and the amino acid sequence deduced therefrom are depicted in SEQ ID NO: 13 therein. FIG. 1 shows a restriction enzyme map of human PGIS cDNA and the region of human PGIS cDNA, which corresponds to the DNA contained in λ hPGIS141, pHPGIS36 and pHPGIS135. FIG. 2 shows a restriction enzyme map of pHPGIS36 and FIG. 3 shows a restriction enzyme map of pHPGIS135.

[124] Human PGIS cDNA obtained by the above-mentioned cloning had a consensus sequence of the initiation codon of eukaryotic shown by Kozak et al [Nucleic Acids Res. 12, 857-872 (1984)] at around the translation initiation codon, and TGA codon corresponding to the termination codon at 500 codons therefrom. Based on these facts, it was found that the

cDNA of the cloned human PGIS comprised 1977 bp comprising 1500 bp encoding 500 amino acid residues, as shown in SEQ ID NO: 15, and the molecular weight of the protein coded thereby was speculated to be about 57,000.

[125] Comparison of the amino acid sequence encoded by said DNA with the amino acid sequence of bovine-originated PGIS separately cloned by the present inventor revealed an about 88% homology. The study of bovine PGIS by the present inventor found that the bovine PGIS had a 31% homology with cholesterol 7 α -hydroxylase belonging to the cytochrome P450 7 family (CYP7), and the region around the 441st Cys residue, which is heme-binding site (fifth ligand) of cytochrome P450, was reserved. The human PGIS similarly reserved the amino acid sequence corresponding to said region, and this region is considered to play an important role in the PGIS activity.

[126] Although the bovine PGIS had a 31% homology with cholesterol 7 α -hydroxylase, it had only a 16% homology with human thromboxane synthase belonging to the cytochrome P450 family and a not more than 40% homology with any of the known cytochrome P450 proteins. It is postulated, therefore, that it is a new family in the cytochrome P450 super family, and human PGIS also belongs to this new family.

[127] A search for such structural correlation in activity is indispensable for the study and development of pharmaceutical products. Such search is accomplished only after the primary structure of human PGIS has been clarified. Accordingly, the present invention which discloses the primary structure of human PGIS for the first time is extremely important and significant for the research, and from industrial aspect as well.

Example 2

Expression of human PGIS

(1) Construction of expression vector for human PGIS

[128] A cDNA insert region is cleaved out respectively from the obtained pHPGIS36 clone and pHPGIS135 clone using a suitable restriction enzyme, and purified. The both fragments obtained were thermally denatured (95°C for 10 minutes), followed by annealing. cDNA is replicated using a DNA polymerase to the both directions toward 5' and 3' from the overlapped region as the synthesis initiation region. Using the obtained whole length cDNA as a template, a primer is synthesized from each region of initiation codon or termination codon and PCR is performed. On this occasion, a suitable restriction enzyme site is constructed as an anchor site at 3' of the primer.

[129] The PCR product thus obtained is purified, the nucleotide sequence of which is confirmed, and the product is digested with BamHI and SmaI (BglII) to give a BamHI-SmaI (BglII) fragment. Said BamHI-SmaI (BglII) fragment is introduced into the BamHI-SmaI site of pVL1393 expression vector previously treated with BamHI-SmaI. The recombinant plasmid thus formed (PGIS7) is characterized by restriction enzyme mapping and DNA sequence analysis.

(2) Baculovirus expression system

[130] Sf9 cells (manufactured by In Vitrogen) are mono-layer cultured in a Grace's insect medium containing 10% fetal calf serum, 0.33% yeastolate and 0.33% lactoalbumin hydrolysate at 27°C. For the production of a recombinant virus, Sf9 cell (1.5×10^6 cells) recombinant plasmid (PGIS7, 50 µg) and wild type baculovirus DNA (AcNPV; 1 µg) are mixed and transfected by calcium phosphate precipitation method. The recombinant baculovirus is isolated and amplified by a combination of plaque assay and slot hybridization using a ^{32}P -labeled cDNA fragment of PGIS as a probe.

[131] Said Sf9 cells are infected with wild type baculovirus or recombinant baculovirus. At 3 days after the infection, cells are collected (2×10^8 cells) and incubated for 5 hours in a serum-containing medium with or without 10 µM hemin.

[132] The obtained cells are washed with phosphate-buffered saline and preserved at -80°C. The microsomal fraction of the cell is prepared according to the method of Haurand and Ullrich et al. [J. Biol. Chem. 260, 15059-15067 (1985)]. The obtained cells (2×10^8 cells) are homogenized in a solution (20 ml) of 10 mM potassium phosphate buffer (pH 7.0), 10 mM EDTA, 5 mM glucose, 0.1 mM dithiothreitol (DTT), 1.15% KCl, 2 µg/ml leupeptin, 2 µg/ml pepstatin, 10 µg/ml soybean trypsin inhibitor and 44 µg/ml phenylmethylsulfonyl fluoride, and subjected to ultrasonication (30 seconds, 4 times) using a Branson sonifier model 450.

[133] The obtained homogenate is centrifuged at 7,000 X g for 15 minutes, and the obtained supernatant is centrifuged at 105,000 X g for 60 minutes. The sediment obtained is suspended in 10 mM potassium phosphate buffer (3 ml, pH 7.0) containing 20% glycerol, 1 mM DTT and 1 mM EDTA by sonication. The protein concentration is determined by Lowry method using bovine serum albumin as a standard, and a solution for immunoblot analysis and PGIS assay at 5 mg/ml is prepared.

[134] (3) Western immunoblot analysis

[135] The infected Sf9 cells and human platelet microsomal fraction are subjected to 10% SDS-PAGE according to the method of Laemmli [Nature 227, 680-685 (1979)]. The migrated protein is electrophoretically transferred onto a polyvinylidene difluoride (PVDF) membrane (Immobilon, Millipore) according to the method of Towbin et al. [Proc. Natl. Acad. Sci. USA 76,4350-4354 (1979)] Tris-HCl buffered saline (TBS) (pH 7.4) containing 10% equine serum is pretreated at room temperature for 30 minutes, and the blot membrane is incubated with polyclonal antibody against bovine PGIS in TBS containing 3% skim milk.

[136] After washing with TBS containing 0.05% Tween 20, the membrane is incubated in TBS containing 3% skim milk at 37°C for 30 minutes together with anti-mouse IgG equine antibody conjugated with horseradish peroxidase (manufactured by Vector Laboratories).

After thorough washing with TBS containing 0.05% Tween 20, the band showing positive immunological response is detected using an immunostaining HRP kit (manufactured by Konica).

[137] Example 3

[138] Expression of human PGIS in cultured animal cell

[139] (1) Preparation of whole length human PGIS cDNA

[140] The obtained pHPGIS36 clone was cleaved out with restriction enzymes SalI and NspI and purified to give a SalI-NspI fragment. The pHPGIS135 clone was cleaved out with restriction enzymes PstI and BamHI and purified to give a PstI-BamHI fragment.

Furthermore, primers [SEQ ID NO: 16: P5 primer (676-699), SEQ ID NO: 17: P6 primer (832-855)] having sequences depicted in Sequence Listing SEQ ID NOS: 16 and 17 were synthesized based on the nucleotide sequence of λ hPGIS141. Using these primers and λ hPGIS141 as a template, a middle stream region of human PGIS cDNA was amplified by PCR method, cleaved with restriction enzymes NspI and PstI, purified and confirmed for the nucleotide sequence and used as an NspI-PstI fragment. These SalI-NspI fragment, PstI-BamHI fragment and NspI-PstI fragment were bound and introduced into the SalI-BamHI site of pBluescriptII SK+ (manufactured by STRATAGENE) previously treated with SalI-BamHI, whereby a plasmid (pHPGIS1) containing the whole length human PGIS cDNA was prepared. FIG. 4 shows the restriction enzyme map of pHPGIS 1.

[141] (2) Construction of human PGIS expression vector for cultured animal cell

[142] Human PGIS cDNA insert region was cleaved out from the obtained pHPGIS1 clone with restriction enzymes SalI and BamHI and purified to give a SalI-BamHI fragment. This SalI-BamHI fragment was introduced into the SalI-BamHI site of pCMV7 expression vector

[supplied by Dr. David W. Russel, University of Texas Southwestern Medical Center, Cell, 75, 187-197 (1993); J. Biol. Chem., 264, 8222-8229 (1989)] previously treated with SalI-BamHI, whereby a human PGIS expression vector (pCMV-HPGIS 1) for cultured animal cell was prepared. FIG. 5 shows the restriction enzyme map of pCMV-HPGIS 1.

[143] (3) Expression of human PGIS in cultured animal cell

[144] Human fetus kidney-derived 293 cells (manufactured by Dainippon Pharmaceutical Co., Ltd.) were sewn in a 60 mm dish at 3×10^5 cells, and mono-layer cultured at 37°C for 24 hours in Dulbecco modified Eagle's medium (DMEM) containing 10% fetal calf serum, 100 U/ml penicillin and 100 µg/ml streptomycin. Then, a recombinant plasmid (pCMV-HPGIS 1, 3 µg) and pVA1 [adenovirus VA1 gene, 3 µg: supplied by Dr. David W. Russel, University of Texas Southwestern Medical Center, Mol. Cell. Biol., 7, 549-551 (1987)] were mixed and transfected by lipofectin method (GIBCO BRL). At 40 hours after the transfection, the cells were washed with phosphate-buffered saline and collected. The cells were suspended in 10 mM calcium phosphate buffer (pH 7.0) containing 10 mM EDTA, 10 mM phenylmethanesulfonyl fluoride (PMSF), 5 mM glucose, 0.1 mM dithiothreitol (DTT), 1.15% KCl, 2 µg/ml leupeptin, 2 µg/ml pepstatin and 10 µg/ml soybean trypsin inhibitor, and subjected to ultrasonication (10 seconds, 10 times) using ASTRASON™ Model XL2020.

[145] The obtained homogenate was centrifuged at 100,000 X g for 60 minutes and the obtained sediment was suspended in 10 mM calcium phosphate buffer (pH 7.0) containing 1 mM EDTA, 1 mM PMSF, 20% glycerol and 0.1 mM DTT. The protein concentration of the obtained sample was determined using a BCA (bicinchoninic acid) protein concentration determination kit (manufactured by PIERCE) using bovine serum albumin as a standard. The PGIS activity of the obtained sample was determined by reacting same with ^{14}C -labeled

PGH₂ (5 nmole) as a substrate at 24°C for 2 minutes, separating 6-keto-PGF₁α, which is a metabolite of the produced PGI₂, by thin layer chromatography, and detecting the radioactivity of the 6-keto-PGF₁α. FIG. 7 shows the detected PGIS activity, FIG. 8 shows PGIS activity of positive control (bovine platelet microsomes) and FIG. 9 shows the analysis results, by thin layer chromatography, of negative control wherein pCMV7 alone was introduced.

[146] As the result of the determination using a sample prepared from the cell into which an expression vector incorporating human PGIS cDNA had been introduced, a spot of 6-keto-PGF₁α, which is a metabolite of PGI₂, was detected as shown by an arrow in FIG. 7. The results were the same as those obtained using bovine platelet microsome containing PGIS as a positive control (FIG. 8). In contrast, the determination using a sample prepared from the cell into which an expression vector without human PGIS cDNA had been introduced failed to detect a spot of 6-keto-PGF₁α. The spot of PGH₂ was thicker (FIG. 9) than in FIG. 7 and FIG. 8. The above results mean that PGIS cDNA incorporated in the expression vector was expressed as a recombinant protein (recombinant PGIS) having PGIS activity and this protein acted on PGH₂ to produce 6-keto-PGF₁α which is a metabolite of PGI₂.

[147] Example 4

[148] Expression of human PGIS in cultured animal cell

[149] Human PGIS cDNA was bound to the XhoI site of the expression vector pUC-CAGGS [having an enhancer of cytomegalovirus and chicken beta-actin promoter] as shown in FIG. 6 [prepared according to the description in Gene 108, 193-200 (1991)] to construct an

expression vector. Two kinds of vectors, i.e., this vector and a vacant vector without human PGIS cDNA, were introduced into vascular smooth muscle cells respectively prepared from rat aorta by HVJ-liposome method [Hypertension 21, 894-899 (1993)] and incubated in a serum-free medium [Dulbecco modified Eagle's medium (DMEM) containing 5×10^{-7} M insulin, 50 $\mu\text{g/ml}$ transferin, 0.2 mM ascorbic acid, 100 U/ml penicillin and 100 $\mu\text{g/ml}$ streptomycin] in a CO_2 incubator at 37°C for 2 days. Then, the medium was changed to a medium containing 1% or 5% fetal calf serum (FCS), and ^3H -thymidine was added 16 hours later. At 8 hours after the addition of thymidine, the thymidine uptake was determined by a conventional method [Cancer Immunol. Immunother. 24, 158-164 (1987)].

[150] The results are shown in FIG. 10, wherein control was a cell into which a vacant vector was introduced and PGIS was a cell into which an expression vector bound with human PGIS cDNA was introduced.

[151] Addition of serum to the vascular smooth muscle cell cultured in the absence of serum led to a promoted proliferation which increased thymidine uptake. In the vascular smooth muscle cell into which an expression vector ligated with human PGIS cDNA was introduced, thymidine intake, namely, proliferation, was significantly suppressed as compared to the cell into which a vacant vector was introduced. This result suggests the possibility of PGIS cDNA introduction suppressing abnormal growth of smooth muscle cells in vascular intima which is observed in arteriosclerosis and the like.

[152] Example 5

[153] Preparation of anti-PGIS polyclonal antibody

[154] PGIS dissolved in 0.5 ml of phosphate-buffered saline (PBS) and an equivalent amount of adjuvant were emulsified and subcutaneously injected to rabbit. Thereafter, similar subcutaneous injection was given twice every 10 days, and blood was taken from the rabbit 10 days after the final subcutaneous injection. Anti-PGIS/IgG was purified and obtained from rabbit anti-PGIS serum prepared from the blood of said rabbit using protein A sepharose 4B (Bio-Rad).

[155] Example 6

[156] Preparation of anti-PGIS monoclonal antibody

[157] (1) Mouse

[158] Male inbred line BALB/c mice (5 weeks of age) were obtained and bred on standard pellet in an animal breeding chamber (23±1°C, 70% humidity) with optional watering.

[159] (2) Immunogen

[160] Human-originated purified PGIS was used. The human PGIS was prepared to a concentration of 1 mg/ml with Dulbecco PBS, dispensed into test tubes by 100 µg and freeze-preserved at -80°C until use.

[161] (3) Immunizing method

[162] Human PGIS 100 µg/0.5 ml and an equivalent amount of Freund's complete adjuvant were mixed. An emulsified antigen (20 µg) was administered to five male BALB/c mice (5 weeks of age) intraperitoneally and subcutaneously at dozen sites on the back every 2 weeks for 2 months. After the immunization for 2 months, antibody titer was measured, and the mice having high antibody titer were picked and applied with additional intraperitoneal administration of 50 µg, 100 µg or 200 µg thereof every other week.

[163] After the immunization for 2 months, two different mice were intraperitoneally administered with 100 µg thereof after a blank of one month. One week later, 100 µg thereof was intravenously injected for additional immunization.

[164] (4) Cell fusion

[165] At 3 days from the final immunization, the spleen of the BALB/c mice was removed to prepare suspensions of spleen cells in EMEM culture medium. The spleen cells were washed 4 times with EMEM culture medium and counted.

[166] For cell fusion, 2-amino-6-oxy-8 azapuraine (8-Azaguanine)-resistant BALB/c mouse myeloma-derived cultured cell line (P3-X63-Ag8.653, hereinafter abbreviated as X63 cells) was used as a parent cell line. The X63 cells were subcultured in RPMI-1640 culture medium (20 µg/ml, containing 8-Azaguanine) supplemented with 5% inactivated fetal calf serum (FCS), and X63 cells in the logarithmic growth phase were washed 3 times with RPMI-1640 culture medium and counted.

[167] Cell fusion is performed in RPMI-1640 culture medium containing polyethylene glycol 4000 at a concentration of 50 (w/v) %.

[168] That is, spleen cells and X63 cells are mixed at a ratio of 10:1 and centrifuged at 1500 rpm for 5 minutes. Supernatant is removed, and cell pellets are thoroughly suspended and subjected to cell fusion according to the method of Kohler and Milstein using polyethylene glycol. Thereafter, the spleen cells are suspended in an HAT selective medium (10% FCS-added RPMI-1640 culture medium containing 1×10^{-4} M hypoxanthine, 4×10^{-7} M aminopterin and 1.6×10^{-5} M thymidine) so that the spleen cells are contained at a concentration of 3.5×10^6 cells/ml. Then, the cell suspension is dispensed into each well of 96 well microtest plate by 100 μ l and cultured in a carbonic acid gas incubator (37°C, 95% humidity, 8% carbonic acid gas). On day 1 and day 2 after the initiation of culture, HAT medium is added by one drop to each well and by 2 drops on day 7 and day 9 after the initiation of incubation, which is followed by further culture.

[169] (5) Screening

[170] From 10 days after the initiation of culture, clone cells emerge. For confirmation of antibody production, hybridoma culture supernatant is subjected to an antigen-antibody reaction test.

[171] That is, 50 μ l each from hybridoma culture supernatant and human PGIS antigen liquid is placed in a U-bottomed microtiter plate and thereto is added 50 μ l of 20% suspension of Sepharose 4B bound with anti-mouse immunoglobulin antibody. The mixture is stirred at room temperature for one hour and left standing for 10 minutes. After confirmation of complete sedimentation of anti-mouse immunoglobulin antibody-bound Sepharose 4B on the bottom of the well, 20 μ l of the supernatant is taken and determined for concentration of residual human PGIS in the supernatant by PGIS ELISA system. When anti-human PGIS monoclonal antibody against human PGIS is present in the hybridoma culture supernatant,

human PGIS and anti-human PGIS monoclonal antibody react and anti-mouse immunoglobulin antibody-bound Sepharose 4B sediment is formed as an antigen-antibody complex to decrease the concentration of residual human PGIS in the supernatant, thus proving the presence of anti-human PGIS monoclonal antibody.

[172] Example 7

[173] A synthetic peptide (with 14 residues), GCGIEALPRTHESQ (SEQ ID NO: 20) corresponding to amino acid residues 182-193 of human PGIS, fused to Gly-Cys at the N-terminus, was coupled to keyhole limpet hemocyanin (KLH) by standard method. Thus, the peptide itself is GIEALPRTHESQ (SEQ ID NO: 18). The corresponding sequence for bovine PGIS is GVEAPPHTQESQ (SEQ ID NO: 19). A Japanese white rabbit was injected with 700 µg of the conjugated peptide in Freund's complete adjuvant subcutaneously, and was boosted every 2 weeks, 3 times (700 µg of the conjugate in Freund's incomplete adjuvant). After one week from the third boost, the blood was collected from the rabbit carotid artery and the serum was used for immunoblotting as antiserum against human PGIS.

[174] The human and bovine PGIS were prepared using the BAC- to BACTM Baculovirus Expression System (Invitrogen). The entire protein coding region of human or bovine cDNA was inserted into the pFastBac-1 donor plasmid. The constructed plasmids were transfected into DH10Bac competent cells, and the recombinant bacmid DNAs and the recombinant baculovirus generated according to the manufacture's instruction. The recombinant baculovirus or empty (Mock) baculovirus were infected to *Spodoptera frugiperda* 21 (Sf21) insect cells and the cells were cultured at 27°C in Grace's insect cell culture medium supplemented with 10% fetal bovine serum, 0.4% yeastolate, 0.4% lactalbumin hydrolysate and 5mg/ml hematin. After 72 h, the cells were harvested and solubilized microsome was

prepared according to J. Biol. Chem., 1994, 269(31):19897-19903. Solubilized microsome proteins (2 µg/lane) were separated by 10% SDS-polyacrylamide gel electrophoresis according to the method of Laemmli. For immunoblotting analysis, proteins were transferred electrophoretically onto a polyvinylidene fluoride membrane (Millipore) for 1.5h at 1.5 mA/cm². The membrane was blocked with 5% skim-milk in 20 mM Tris-HCl, pH 7.5 containing 500 mM NaCl and 0.05% Tween 20 (Tris-buffered salt solution with Tween 20, TBS-T) at room temperature overnight. Subsequently, the membrane was blocked with 1% BSA in TBS-T at room temperature for 1 h and then incubated with the antiserum (x 200) against human PGIS for 1 h. After the membrane was rinsed with TBS-T 3 times for 10 min, it was incubated with horseradish peroxidase conjugated anti-rabbit IgG (x1000, Daco) in TBS-T containing 1% BSA for 1 h. After washing with TBS-T 4 times for 10 min., the membrane was reacted with the ECL detection reagents (Amersham Pharmacia Biotech Ltd.) and exposed to X-ray film. For stripping antibodies from the membrane, the membrane was washed with 62.5 mM Tris-HCl 6.7 containing 2% SDS and 0.1 M 2-mercaptoethanol at 50°C for 30 min and then in TBS-T at room temperature twice for 10 min. After blocking, the membrane was reacted with the polyclonal antiserum (x1000) against murine PGIS (whole) and the secondary antibody, and then the signal was detected as described above. The results are shown in Figure 14.

[175] Figure 14 displays the detected signals on the membrane reacted with the antiserum against human (left panel) or murine (right panel) PGIS. Lane 1, human PGIS-infected Sf21 cells; lane 2, bovine PGIS-infected Sf21 cells, lane 3, Mock-infected Sf21 cells.

[176] From the above experiment, it was found that antiserum against human PGIS reacted with human PGIS, but did not react with bovine PGIS. On the other hand, antiserum against murine PGIS cross-reacted with both human and bovine PGIS. These results clearly demonstrate that the inventive antibody is highly specific to human PGIS.

[177] The human “synthetic” peptide used in this Example 7 is SEQ ID NO: 18 and the corresponding bovine PGIS sequence is SEQ ID NO: 19, differing from the human PGIS sequence at four amino acid locations.

[178] Example 8

[179] Purification of anti-PIS polyclonal antibody

[180] The anti-PGIS polyclonal antibody, prepared in Example 7, is purified by well known techniques, such as ammonium sulfate precipitation, gel filtration chromatography, ion exchange chromatography or affinity chromatography using protein A or protein G, which provide primarily the IgG fraction of immune serum. For the further purification of the crude anti-PGIS IgG fraction by affinity chromatography, the synthetic peptide with the sequence GIEALPRTHESQ is covalently immobilized on a Sepharose. Crude anti-PGIS IgG fraction is applied to the peptide-conjugated- Sepharose, and subsequently washed with PBS. The antibodies are eluted from the peptide-conjugated-Sepharose with 0.1 mol/l acetic acid (pH 2.5), and the antibody solution is neutralized with solid sodium phosphate (0.01 mol/l).

[181] Reference Example 1

[182] RNA blot analysis

[183] RNA blot hybridization analysis was made to examine the influence of several kinds of cytokines on the expression of HAEC-derived human PGIS mRNA.

[184] The entire RNA (30 µg) derived from each HAEC which was incubated for 24 hours with several kinds of cytokines [IL-1 α (1 ng/ml), IL-1 β (1 ng/ml), IL-6 (2.5 ng/ml), TNF- α (5 ng/ml) and TNF- β (1 ng/ml)] was denatured with formamide, electrophoresed on 1% agar gel containing 1.5% formaldehyde, and transferred onto a nylon membrane. A probe

[pHPGIS 135 and glyceraldehyde-3-phosphate dehydrogenase (GAPDH)] was labeled with [α -³²P]dCTP by random priming method [Feinberg, A. P., and Vogelstein, B. (1983) *Anal. Biochem.* 132, 6-13].

[185] Then, hybridization was applied according to the method described in *Biochem. Biophys. Res. Commun.* 178, p 1479-1484 (1991). The membrane obtained was washed with 0.1 X SSC (0.15M NaCl, 0.015M sodium citrate, pH 7.0) containing 0.1% SDS at 60° C, air-dried and autoradiographed. The results are shown in FIG. 11. The main band of the HAEC-derived human PGIS mRNA was found at about 6 kb and three other minor bands were found (3.2, 2.5 and 1.7 kb). The test results revealed that the expression of human PGIS mRNA incubated for 24 hours with IL-1 α , IL-1 β or IL-6 increased about 2-fold as compared with the control without cytokine treatment. Accordingly, increase in PGI₂ production caused by cytokine is considered to be attributable to the increased expression and production of PGIS which was achieved by cytokine. Thus, the treatment with cytokine is an extremely useful method for increasing PGIS expression to increase PGIS activity, which in turn accelerates PGI₂ production.

[186] Reference Example 2

[187] In vivo distribution of PGIS mRNA

[188] RNA blot analysis was made to examine the distribution of PGIS mRNA expression in human body. Specifically, a filter was purchased from Clone-Tech on which poly (A)⁺ RNA of various human tissues was electrophoresed and blotted. hPGIS135 was labeled with ³²P by the aforementioned method and subjected to Northern blot hybridization under the same conditions as above.

[189] The results are shown in FIG. 12 and FIG. 13. The results confirm that PGIS mRNA was abundantly expressed widely in human tissues, particularly, in uterus, heart, skeletal muscle, lung and prostate and at significant levels, though slightly, in small intestine, kidney, liver and brain. These results coincide with the conventional reports of enzymatic activity and distribution in tissue of immunological response of PGIS, thus suggesting various biological roles assumed by PGIS besides the action in the vascular system. The 6 kb main, strong band and 3 weak bands as shown in FIG. 11 were observed in all tissues mentioned above, though relative thickness among the weak bands varied between tissues. Such various modes of presence of transcription products suggest possible different splicing of mRNA or the presence of an analogous gene (isozyme) as found in prostaglandin endoperoxidase.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(iii) NUMBER OF SEQUENCES: 19

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 16 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GGGACAAGGACCACAT16

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

CAAAAGTCGCCTGTGGAAGC20

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 18 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

CACAGGCGACTTTTGACA18

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 18 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

TGCCTGCATCTCCTCTGA18

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 35 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: other nucleic acid

(A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

GACTCGAGTCGACATCGATTTTTTTTTTTTTTTTTTTT35

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 17 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: other nucleic acid
 (A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
 GACTCGAGTCGACATCG17
 (2) INFORMATION FOR SEQ ID NO:7:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 48 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: other nucleic acid
 (A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC RNA"
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
 CUACUACUACUAGGCCACGCGTCGACTAGTACGGGNNGGGNNGGGNN48
 (2) INFORMATION FOR SEQ ID NO:8:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 183 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: cDNA
 (vi) ORIGINAL SOURCE:
 (A) ORGANISM: Homo sapiens
 (B) STRAIN: λ hPGIS141
 (ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 1..183
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
 GGGGACAAGGACCACATGTGCAGTGTCAAAGTCGCCTGTGGAAGCTG48
 GlyAspLysAspHisMetCysSerValLysSerArgLeuTrpLysLeu
 151015
 CTATCCCCAGCCAGGCTGGCCAGGCGGGCCCCACCGGAGCAAATGGCTG96
 LeuSerProAlaArgLeuAlaArgArgAlaHisArgSerLysTrpLeu
 202530
 GAGAGTTACCTGCTGCACCTGGAGGAGATGGGTGTGTCAGAGGAGATG144
 GluSerTyrLeuLeuHisLeuGluGluMetGlyValSerGluGluMet
 354045
 CAGGCACGGGCCCTGGTGTGCTGCAGCTGTGGGCCACACAG183
 GlnAlaArgAlaLeuValLeuGlnLeuTrpAlaThrGln
 505560
 (2) INFORMATION FOR SEQ ID NO:9:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 61 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: protein
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
 GlyAspLysAspHisMetCysSerValLysSerArgLeuTrpLysLeu
 151015
 LeuSerProAlaArgLeuAlaArgArgAlaHisArgSerLysTrpLeu
 202530
 GluSerTyrLeuLeuHisLeuGluGluMetGlyValSerGluGluMet
 354045
 GlnAlaArgAlaLeuValLeuGlnLeuTrpAlaThrGln
 505560
 (2) INFORMATION FOR SEQ ID NO:10:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 792 base pairs

(B) TYPE: nucleic acid
 (C) STRANDEDNESS: double
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: cDNA
 (vi) ORIGINAL SOURCE:
 (A) ORGANISM: Homo sapiens
 (B) STRAIN: pHPGIS36
 (ix) FEATURE:
 (A) NAME/KEY: CDS
 (B) LOCATION: 80..790
 (ix) FEATURE:
 (A) NAME/KEY: mat.sub.-- peptide
 (B) LOCATION: 80..790
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
 CTACTACTACTAGGCCACGCGTCGACTAGTACGGGGGGGGGGGGGGGGGGCAGCCCCGC60
 CAGCCCCGCCAGCCCCGCATGGCTTGGGCCGCGCTCCTCGGCCTCCTGGCC112
 MetAlaTrpAlaAlaLeuLeuGlyLeuLeuAla
 1510
 GCACTGTTGCTGCTGCTGCTACTGAGCCGCCGCCGACGCGGCGACCT160
 AlaLeuLeuLeuLeuLeuLeuSerArgArgArgThrArgArgPro
 152025
 GGTGAGCCTCCCCGACCTGGGCAGCATCCCCGTTGGTGGGGTATGCC208
 GlyGluProProLeuAspLeuGlySerIleProTrpLeuGlyTyrAla
 303540
 TTGGACTTTGGAAAAGATGCTGCCAGCTTCCTCAGGAGATGAAGGAG256
 LeuAspPheGlyLysAspAlaAlaSerPheLeuThrArgMetLysGlu
 455055
 AAGCACGGTGACATCTTTACTATACTGGTTGGGGGCAGGTATGTCACC304
 LysHisGlyAspIlePheThrIleLeuValGlyGlyArgTyrValThr
 60657075
 GTTCTCCTGGACCCACACTCCTACGACGCGGTGGTGTGGGAGCCTCGC352
 ValLeuLeuAspProHisSerTyrAspAlaValValTrpGluProArg
 808590
 ACCAGGCTCGACTTCCATGCCTATGCCATCTTCCTCATGGAGAGGATT400
 ThrArgLeuAspPheHisAlaTyrAlaIlePheLeuMetGluArgIle
 95100105
 TTTGATGTGCAGCTTCCACATTACAGCCCCAGTGATGAAAAGGCCAGG448
 PheAspValGlnLeuProHisTyrSerProSerAspGluLysAlaArg
 110115120
 ATGAACTGACTCTTCTCCACAGAGAGCTCCAGGCACTCACAGAAGCC496
 MetLysLeuThrLeuLeuHisArgGluLeuGlnAlaLeuThrGluAla
 125130135
 ATGTATACCAACCTCCATGCAGTGCTGTTGGGCGATGCTACAGAAGCA544
 MetTyrThrAsnLeuHisAlaValLeuLeuGlyAspAlaThrGluAla
 140145150155
 GGCAGTGGCTGGCACGAGATGGGTCTCCTCGACTTCTCCTACAGCTTC592
 GlySerGlyTrpHisGluMetGlyLeuLeuAspPheSerTyrSerPhe
 160165170
 CTGCTCAGAGCCGGCTACCTGACTCTTTACGGAATTGAGGCGCTGCCA640
 LeuLeuArgAlaGlyTyrLeuThrLeuTyrGlyIleGluAlaLeuPro
 175180185
 CGCACCCATGAAAGCCAGGCCAGGACCGCTCCACTCAGCTGATGTC688
 ArgThrHisGluSerGlnAlaGlnAspArgValHisSerAlaAspVal
 190195200
 TTCCACACCTTTGCGCCAGCTCGACCGGCTGCTCCCCAACTGGCCCGT736
 PheHisThrPheArgGlnLeuAspArgLeuLeuProLysLeuAlaArg
 205210215
 GGCTCCCTGTCACTGGGGGACAAGGACCACATGTGCAGTGTCAAAGT784
 GlySerLeuSerValGlyAspLysAspHisMetCysSerValLysSer
 220225230235

CGCCTGTG792

ArgLeu

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 237 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

MetAlaTrpAlaAlaLeuLeuGlyLeuLeuAlaAlaLeuLeuLeuLeu
151015

LeuLeuLeuSerArgArgArgThrArgArgProGlyGluProProLeu
202530

AspLeuGlySerIleProTrpLeuGlyTyrAlaLeuAspPheGlyLys
354045

AspAlaAlaSerPheLeuThrArgMetLysGluLysHisGlyAspIle
505560

PheThrIleLeuValGlyGlyArgTyrValThrValLeuLeuAspPro
65707580

HisSerTyrAspAlaValValTrpGluProArgThrArgLeuAspPhe
859095

HisAlaTyrAlaIlePheLeuMetGluArgIlePheAspValGlnLeu
100105110

ProHisTyrSerProSerAspGluLysAlaArgMetLysLeuThrLeu
115120125

LeuHisArgGluLeuGlnAlaLeuThrGluAlaMetTyrThrAsnLeu
130135140

HisAlaValLeuLeuGlyAspAlaThrGluAlaGlySerGlyTrpHis
145150155160

GluMetGlyLeuLeuAspPheSerTyrSerPheLeuLeuArgAlaGly
165170175

TyrLeuThrLeuTyrGlyIleGluAlaLeuProArgThrHisGluSer
180185190

GlnAlaGlnAspArgValHisSerAlaAspValPheHisThrPheArg
195200205

GlnLeuAspArgLeuLeuProLysLeuAlaArgGlySerLeuSerVal
210215220

GlyAspLysAspHisMetCysSerValLysSerArgLeu
225230235

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1296 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Homo sapiens

(B) STRAIN: pHPGIS135

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 3..827

(ix) FEATURE:

(A) NAME/KEY: mat.sub.-- peptide

(B) LOCATION: 3..827

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

GGGACAAGGACCACATGTGCAGTGTCAAAGTCGCCTGTGGAAGCTG47

AspLysAspHisMetCysSerValLysSerArgLeuTrpLysLeu

151015

CTATCCCCAGCCAGGCTGGCCAGGCGGGCCCCACCGGAGCAAATGGCTG95

LeuSerProAlaArgLeuAlaArgArgAlaHisArgSerLysTrpLeu
 202530
 GAGAGTTACCTGCTGCACCTGGAGGAGATGGGTGTGTCAGAGGAGATG143
 GluSerTyrLeuLeuHisLeuGluGluMetGlyValSerGluGluMet
 354045
 CAGGCACGGGCCCTGGTGTCTGCAGCTGTGGGCCACACAGGGGAATATG191
 GlnAlaArgAlaLeuValLeuGlnLeuTrpAlaThrGlnGlyAsnMet
 505560
 GGTCCCGCTGCCTTCTGGCTCCTGCTCTTCCTTCTCAAGAATCCTGAA239
 GlyProAlaAlaPheTrpLeuLeuLeuPheLeuLeuLysAsnProGlu
 657075
 GCCCTGGCTGTGTCCGCGGAGAGCTCGAGAGTATCCTTTGGCAAGCG287
 AlaLeuAlaAlaValArgGlyGluLeuGluSerIleLeuTrpGlnAla
 80859095
 GAGCAGCCTGTCTCGCAGACGACCACTCTCCACAGAAGGTTCTAGAC335
 GluGlnProValSerGlnThrThrThrLeuProGlnLysValLeuAsp
 100105110

AGCACACCTGTGCTTGATAGCGTGCTGAGTGAGAGCCTCAGGCTTACA383
 SerThrProValLeuAspSerValLeuSerGluSerLeuArgLeuThr
 115120125
 GCTGCCCCCTTCATCACCCGCGAGGTTGTGGTGGACCTGGCCATGCCC431
 AlaAlaProPheIleThrArgGluValValValAspLeuAlaMetPro
 130135140
 ATGGCAGACGGGAGAGAATTCAACCTGCGACGTGGTGACCGCCTCCTC479
 MetAlaAspGlyArgGluPheAsnLeuArgArgGlyAspArgLeuLeu
 145150155
 CTCTTCCCCTTCTGAGCCCCAGAGAGACCCAGAAATCTACACAGAC527
 LeuPheProPheLeuSerProGlnArgAspProGluIleTyrThrAsp
 160165170175
 CCAGAGGTATTTAAATACAACCGATTCTGAACCTGACGGATCAGAG575
 ProGluValPheLysTyrAsnArgPheLeuAsnProAspGlySerGlu
 180185190
 AAGAAAGACTTTTACAAGGATGGGAAACGGCTGAAGAATTACAACATG623
 LysLysAspPheTyrLysAspGlyLysArgLeuLysAsnTyrAsnMet
 195200205
 CCCTGGGGGGCGGGGCACAATCACTGCCTGGGGAGGAGTTATGCGGTC671
 ProTrpGlyAlaGlyHisAsnHisCysLeuGlyArgSerTyrAlaVal
 210215220
 AACAGCATCAAACAATTTGTGTTCTTGTGCTGGTGCACCTTGGACTTG719
 AsnSerIleLysGlnPheValPheLeuValLeuValHisLeuAspLeu
 225230235
 GAGCTGATCAACGCAGATGTGGAGATCCCTGAGTTTGACCTCAGCAGG767
 GluLeuIleAsnAlaAspValGluIleProGluPheAspLeuSerArg
 240245250255
 TACGGCTTCGGTCTGATGCAGCCGGAACACGACGTGCCCCGTCGCTAC815
 TyrGlyPheGlyLeuMetGlnProGluHisAspValProValArgTyr
 260265270
 CGCATCCGCCCATGACACAGGGAGCAGATGGATCCACGTGCTCGCCTCTGCC867
 ArgIleArgPro
 275
 CAGCCTGCCCCAGCCTGCCCCAGCCTCCCAGCTTTCTGTGTGCACAGTTGGCCCGGGTGC927
 AGGTGCTAGCATTACCACTTCCCTGCTTTTCTCCCAGAAGGCTGGGTCCAGGGGAGGGAA987
 AAGCTAAGAGGGTGAACAAAGAAAAGACATTGAAAGCTCTATGGATTATCCACTGCAAAG1047
 TTTTCTTTCCAAAATCAGGCTTTGTCTGCTCCCAATTACCTCGTTACTCTCACCTCGTG1107
 ATATCCACAAATGCTATTCAGATAAGGCAGAACTAGGAGTCTTCACTGCTCTGCCCCAA1167
 CTCCCGGAGGTGTACCTTCCTAGTTCTTATGAGCTAGCATGGCCCGGGCCTTATCCAGT1227
 CAAAGCGGATGCTGGCCACAGAAAGGCCACTCAGGATGTCCTTTGTGTCCATCGATGTGCG1287
 ACTCGAGTC1296

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 275 amino acids

(B) TYPE: amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

AspLysAspHisMetCysSerValLysSerArgLeuTrpLysLeuLeu
151015
SerProAlaArgLeuAlaArgArgAlaHisArgSerLysTrpLeuGlu
202530
SerTyrLeuLeuHisLeuGluGluMetGlyValSerGluGluMetGln
354045
AlaArgAlaLeuValLeuGlnLeuTrpAlaThrGlnGlyAsnMetGly
505560
ProAlaAlaPheTrpLeuLeuLeuPheLeuLeuLysAsnProGluAla
65707580
LeuAlaAlaValArgGlyGluLeuGluSerIleLeuTrpGlnAlaGlu
859095
GlnProValSerGlnThrThrThrLeuProGlnLysValLeuAspSer
100105110
ThrProValLeuAspSerValLeuSerGluSerLeuArgLeuThrAla
115120125
AlaProPheIleThrArgGluValValValAspLeuAlaMetProMet
130135140
AlaAspGlyArgGluPheAsnLeuArgArgGlyAspArgLeuLeuLeu
145150155160
PheProPheLeuSerProGlnArgAspProGluIleTyrThrAspPro
165170175
GluValPheLysTyrAsnArgPheLeuAsnProAspGlySerGluLys
180185190
LysAspPheTyrLysAspGlyLysArgLeuLysAsnTyrAsnMetPro
195200205
TrpGlyAlaGlyHisAsnHisCysLeuGlyArgSerTyrAlaValAsn
210215220
SerIleLysGlnPheValPheLeuValLeuValHisLeuAspLeuGlu
225230235240
LeuIleAsnAlaAspValGluIleProGluPheAspLeuSerArgTyr
245250255
GlyPheGlyLeuMetGlnProGluHisAspValProValArgTyrArg
260265270
IleArgPro
275

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1977 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: double

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Homo sapiens

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 28..1527

(ix) FEATURE:

(A) NAME/KEY: mat.sub.-- peptide

(B) LOCATION: 28..1527

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

AGCCCCGCCAGCCCCGCCAGCCCCGCGATGGCTTGGGCCGCGCTCCTCGGC51

MetAlaTrpAlaAlaLeuLeuGly
 15
 CTCCTGGCCGCTGTTGCTGCTGCTGCTACTGAGCCGCCGCCGACG99
 LeuLeuAlaAlaLeuLeuLeuLeuLeuLeuSerArgArgArgThr
 101520
 CGGCGACCTGGTGAGCCTCCCCCTGGACCTGGGCAGCATCCCCCTGGTTG147
 ArgArgProGlyGluProProLeuAspLeuGlySerIleProTrpLeu
 25303540
 GGGTATGCCTTGGACTTTGGAAAAGATGCTGCCAGCTTCCTCACGAGG195
 GlyTyrAlaLeuAspPheGlyLysAspAlaAlaSerPheLeuThrArg
 455055
 ATGAAGGAGAAGCACGGTGACATCTTTACTATACTGGTTGGGGGCAGG243
 MetLysGluLysHisGlyAspIlePheThrIleLeuValGlyGlyArg
 606570
 TATGTCACCGTTCTCCTGGACCCACACTCCTACGACGCGGTGGTGTGG291
 TyrValThrValLeuLeuAspProHisSerTyrAspAlaValValTrp
 758085
 GAGCCTCGCACCAGGCTCGACTTCCATGCCTATGCCATCTTCCTCATG339
 GluProArgThrArgLeuAspPheHisAlaTyrAlaIlePheLeuMet
 9095100
 GAGAGGATTTTTGATGTGCAGCTTCCACATTACAGCCCCAGTGATGAA387
 GluArgIlePheAspValGlnLeuProHisTyrSerProSerAspGlu
 105110115120
 AAGGCCAGGATGAACTGACTCTTCTCCACAGAGAGCTCCAGGCACTC435
 LysAlaArgMetLysLeuThrLeuLeuHisArgGluLeuGlnAlaLeu
 125130135
 ACAGAAGCCATGTATACCAACCTCCATGCAGTGCTGTTGGGCGATGCT483
 ThrGluAlaMetTyrThrAsnLeuHisAlaValLeuLeuGlyAspAla
 140145150
 ACAGAAGCAGGCAGTGGCTGGCAGAGATGGGTCTCCTCGACTTCTCC531
 ThrGluAlaGlySerGlyTrpHisGluMetGlyLeuLeuAspPheSer
 155160165
 TACAGCTTCCTGCTCAGAGCCGGCTACCTGACTCTTTACGGAATTGAG579
 TyrSerPheLeuLeuArgAlaGlyTyrLeuThrLeuTyrGlyIleGlu
 170175180
 GCGCTGCCACGCACCCATGAAAGCCAGGCCAGGACCGCGTCCACTCA627
 AlaLeuProArgThrHisGluSerGlnAlaGlnAspArgValHisSer
 185190195200
 GCTGATGTCTTCCACACCTTTCCGCCAGCTCGACCGGCTGCTCCCCAAA675
 AlaAspValPheHisThrPheArgGlnLeuAspArgLeuLeuProLys
 205210215
 CTGGCCCGTGGCTCCCTGTCAGTGGGGACAAGGACCACATGTGCAGT723
 LeuAlaArgGlySerLeuSerValGlyAspLysAspHisMetCysSer
 220225230
 GTCAAAAGTCGCCTGTGGAAGCTGCTATCCCCAGCCAGGCTGGCCAGG771
 ValLysSerArgLeuTrpLysLeuLeuSerProAlaArgLeuAlaArg
 235240245
 CGGGCCACCGGAGCAAATGGCTGGAGAGTTACCTGCTGCACCTGGAG819
 ArgAlaHisArgSerLysTrpLeuGluSerTyrLeuLeuHisLeuGlu
 250255260
 GAGATGGGTGTGTCAGAGGAGATGCAGGCACGGGCCCTGGTGTGCAG867
 GluMetGlyValSerGluGluMetGlnAlaArgAlaLeuValLeuGln
 265270275280
 CTGTGGGCCACACAGGGGAATATGGGTCCCGCTGCCTTCTGGCTCCTG915
 LeuTrpAlaThrGlnGlyAsnMetGlyProAlaAlaPheTrpLeuLeu
 285290295
 CTCTTCCTTCTCAAGAATCCTGAAGCCCTGGCTGCTGTCCGCGGAGAG963
 LeuPheLeuLeuLysAsnProGluAlaLeuAlaAlaValArgGlyGlu
 300305310
 CTCGAGAGTATCCTTTGGCAAGCGGAGCAGCCTGTCTCGCAGACGACC1011

LeuGluSerIleLeuTrpGlnAlaGluGlnProValSerGlnThrThr
 315320325
 ACTCTCCACAGAAGGTTCTAGACAGCACACCTGTGCTTGATAGCGTG1059
 ThrLeuProGlnLysValLeuAspSerThrProValLeuAspSerVal
 330335340
 CTGAGTGAGAGCCTCAGGCTTACAGCTGCCCCCTTCATCACCCGCGAG1107
 LeuSerGluSerLeuArgLeuThrAlaAlaProPheIleThrArgGlu
 345350355360
 GTTGTGGTGGACCTGGCCATGCCCATGGCAGACGGGAGAGAATTCAAC1155
 ValValValAspLeuAlaMetProMetAlaAspGlyArgGluPheAsn
 365370375
 CTGCGACGTGGTGACCGCTCCTCCTCTTCCCCCTTCCTGAGCCCCCAG1203
 LeuArgArgGlyAspArgLeuLeuLeuPheProPheLeuSerProGln
 380385390
 AGAGACCCAGAAATCTACACAGACCCAGAGGTATTTAAATACAACCGA1251
 ArgAspProGluIleTyrThrAspProGluValPheLysTyrAsnArg
 395400405
 TTCCTGAACCCTGACGGATCAGAGAAGAAAGACTTTTACAAGGATGGG1299
 PheLeuAsnProAspGlySerGluLysLysAspPheTyrLysAspGly
 410415420
 AAACGGCTGAAGAATTACAACATGCCCTGGGGGGCGGGGCACAATCAC1347
 LysArgLeuLysAsnTyrAsnMetProTrpGlyAlaGlyHisAsnHis
 425430435440
 TGCCTGGGGAGGAGTTATGCGGTCAACAGCATCAAACAATTTGTGTTC1395
 CysLeuGlyArgSerTyrAlaValAsnSerIleLysGlnPheValPhe
 445450455
 CTTGTGCTGGTGCACCTTGGACTTGGAGCTGATCAACGCAGATGTGGAG1443
 LeuValLeuValHisLeuAspLeuGluLeuIleAsnAlaAspValGlu
 460465470
 ATCCCTGAGTTTGACCTCAGCAGGTACGGCTTCGGTCTGATGCAGCCG1491
 IleProGluPheAspLeuSerArgTyrGlyPheGlyLeuMetGlnPro
 475480485
 GAACACGACGTGCCCGTCCGCTACCGCATCCGCCCATGACACAGGG1537
 GluHisAspValProValArgTyrArgIleArgPro
 490495500
 AGCAGATGGATCCACGTGCTCGCCTCTGCCAGCCTGCCCCAGCCTGCCCCAGCCTCCCA1597
 GCTTTCTGTGTGCACAGTTGGCCCGGGTGCAGGTGCTAGCATTACCACTTCCCTGCTTTT1657
 CTCCCAGAAGGCTGGGTCCAGGGGAGGGAAAAGCTAAGAGGGTGAACAAAGAAAAGACAT1717
 TGAAAGCTCTATGGATTATCCACTGCAAAGTTTTCTTTCCAAAATCAGGCTTTGTCTGCT1777
 CCCAATTCACCTCGTTACTCTCACCTCGTGATATCCACAAATGCTATTTCAGATAAGGCAG1837
 AACTAGGAGTCTTCACTGCTCTGCCCCCACTCCCGGAGGTGTCACCTTCCTAGTTCTTA1897
 TGAGCTAGCATGGCCCCGGGCCCTTATCCAGTCAAAGCGGATGCTGGCCACAGAAAGGCCAC1957
 TCAGGATGTCCTTTGTGTCC1977
 (2) INFORMATION FOR SEQ ID NO:15:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 500 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: protein
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:
 MetAlaTrpAlaAlaLeuLeuGlyLeuLeuAlaAlaLeuLeuLeuLeu
 151015
 LeuLeuLeuSerArgArgArgThrArgArgProGlyGluProProLeu
 202530
 AspLeuGlySerIleProTrpLeuGlyTyrAlaLeuAspPheGlyLys
 354045
 AspAlaAlaSerPheLeuThrArgMetLysGluLysHisGlyAspIle
 505560
 PheThrIleLeuValGlyGlyArgTyrValThrValLeuLeuAspPro
 65707580

HisSerTyrAspAlaValValTrpGluProArgThrArgLeuAspPhe
 859095
 HisAlaTyrAlaIlePheLeuMetGluArgIlePheAspValGlnLeu
 100105110
 ProHisTyrSerProSerAspGluLysAlaArgMetLysLeuThrLeu
 115120125
 LeuHisArgGluLeuGlnAlaLeuThrGluAlaMetTyrThrAsnLeu
 130135140
 HisAlaValLeuLeuGlyAspAlaThrGluAlaGlySerGlyTrpHis
 145150155160
 GluMetGlyLeuLeuAspPheSerTyrSerPheLeuLeuArgAlaGly
 165170175
 TyrLeuThrLeuTyrGlyIleGluAlaLeuProArgThrHisGluSer
 180185190
 GlnAlaGlnAspArgValHisSerAlaAspValPheHisThrPheArg
 195200205
 GlnLeuAspArgLeuLeuProLysLeuAlaArgGlySerLeuSerVal
 210215220
 GlyAspLysAspHisMetCysSerValLysSerArgLeuTrpLysLeu
 225230235240
 LeuSerProAlaArgLeuAlaArgArgAlaHisArgSerLysTrpLeu
 245250255
 GluSerTyrLeuLeuHisLeuGluGluMetGlyValSerGluGluMet
 260265270
 GlnAlaArgAlaLeuValLeuGlnLeuTrpAlaThrGlnGlyAsnMet
 275280285
 GlyProAlaAlaPheTrpLeuLeuLeuPheLeuLeuLysAsnProGlu
 290295300
 AlaLeuAlaAlaValArgGlyGluLeuGluSerIleLeuTrpGlnAla
 305310315320
 GluGlnProValSerGlnThrThrThrLeuProGlnLysValLeuAsp
 325330335
 SerThrProValLeuAspSerValLeuSerGluSerLeuArgLeuThr
 340345350

AlaAlaProPheIleThrArgGluValValValAspLeuAlaMetPro
 355360365
 MetAlaAspGlyArgGluPheAsnLeuArgArgGlyAspArgLeuLeu
 370375380
 LeuPheProPheLeuSerProGlnArgAspProGluIleTyrThrAsp
 385390395400
 ProGluValPheLysTyrAsnArgPheLeuAsnProAspGlySerGlu
 405410415
 LysLysAspPheTyrLysAspGlyLysArgLeuLysAsnTyrAsnMet
 420425430
 ProTrpGlyAlaGlyHisAsnHisCysLeuGlyArgSerTyrAlaVal
 435440445
 AsnSerIleLysGlnPheValPheLeuValLeuValHisLeuAspLeu
 450455460
 GluLeuIleAsnAlaAspValGluIleProGluPheAspLeuSerArg
 465470475480
 TyrGlyPheGlyLeuMetGlnProGluHisAspValProValArgTyr
 485490495
 ArgIleArgPro
 500

(2) INFORMATION FOR SEQ ID NO:16:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 24 base pairs
 (B) TYPE: nucleic acid

(C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: other nucleic acid
 (A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:
 GACAAGGACCACATGTGCAGTGTC24
 (2) INFORMATION FOR SEQ ID NO:17:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 24 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
 (ii) MOLECULE TYPE: other nucleic acid
 (A) DESCRIPTION: /desc = "PRIMER/SYNTHETIC DNA"
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:
 CTGTGTGGCCACAGCTGCAGCAC24
 (2) INFORMATION FOR SEQ ID NO: 18:
 (i) SEQUENCE CHARACTERISTICS
 (A) LENGTH: 12 AMINO ACIDS
 (B) TYPE: AMINO ACID
 (D) TOPOLOGY: LINEAR
 (ii) MOLECULE TYPE: PROTEIN
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
 GlyIleGluAlaLeuProArgThrHisGluSerGln
 (2) INFORMATION FOR SEQ ID: NO: 19:
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 12 AMINO ACIDS
 (B) TYPE: AMINO ACID
 (D) TOPOLOGY: LINEAR
 (ii) MOLECULE TYPE: PROTEIN
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
 GlyValGluAlaProProHisThrGlnGluSerGln
 (2) INFORMATION FOR SEQ ID NO: 20
 (i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 14 AMINO ACIDS
 (B) TYPE: AMINO ACID
 (D) TOPOLOGY: LINEAR
 (ii) MOLECULE TYPE: PROTEIN
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
 GluCysGlyIleGluAlaLeuProArgThrHisGluSerGln